

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/774,271  
Source: IFW0  
Date Processed by STIC: 09/07/2006

# ***ENTERED***



IFW16

## RAW SEQUENCE LISTING

DATE: 09/07/2006

PATENT APPLICATION: US/10/774,271

TIME: 14:51:04

Input Set : A:\10-774271 - 099425-0100.ST25.txt

Output Set: N:\CRF4\09062006\J774271.raw

3 &lt;110&gt; APPLICANT: Lipton, James M.

5 &lt;120&gt; TITLE OF INVENTION: METHODS AND COMPOUNDS FOR TREATING MALABSORPTION DISEASES

AND

6 INFLAMMATORY CONDITIONS OF THE GASTROINTESTINAL TRACT

8 &lt;130&gt; FILE REFERENCE: 099425-0100

10 &lt;140&gt; CURRENT APPLICATION NUMBER: 10/774,271

11 &lt;141&gt; CURRENT FILING DATE: 2004-02-06

13 &lt;150&gt; PRIOR APPLICATION NUMBER: 60/445,542

14 &lt;151&gt; PRIOR FILING DATE: 2003-02-06

16 &lt;160&gt; NUMBER OF SEQ ID NOS: 19

18 &lt;170&gt; SOFTWARE: PatentIn version 3.3

20 &lt;210&gt; SEQ ID NO: 1

21 &lt;211&gt; LENGTH: 13

22 &lt;212&gt; TYPE: PRT

23 &lt;213&gt; ORGANISM: Homo sapiens

25 &lt;400&gt; SEQUENCE: 1

27 Ser Tyr Ser Met Glu His Phe Arg Trp Gly Lys Pro Val

28 1 5 10

31 &lt;210&gt; SEQ ID NO: 2

32 &lt;211&gt; LENGTH: 3

33 &lt;212&gt; TYPE: PRT

34 &lt;213&gt; ORGANISM: Artificial

36 &lt;220&gt; FEATURE:

37 &lt;223&gt; OTHER INFORMATION: Fragment of SEQ ID NO:1

39 &lt;400&gt; SEQUENCE: 2

41 Lys Pro Val

42 1

45 &lt;210&gt; SEQ ID NO: 3

46 &lt;211&gt; LENGTH: 10

47 &lt;212&gt; TYPE: PRT

48 &lt;213&gt; ORGANISM: Artificial

50 &lt;220&gt; FEATURE:

51 &lt;223&gt; OTHER INFORMATION: Fragment of SEQ ID NO:1

53 &lt;400&gt; SEQUENCE: 3

55 Met Glu His Phe Arg Trp Gly Lys Pro Val

56 1 5 10

59 &lt;210&gt; SEQ ID NO: 4

60 &lt;211&gt; LENGTH: 8

61 &lt;212&gt; TYPE: PRT

62 &lt;213&gt; ORGANISM: Artificial

64 &lt;220&gt; FEATURE:

65 &lt;223&gt; OTHER INFORMATION: Fragment of SEQ ID NO:1

67 &lt;400&gt; SEQUENCE: 4

69 His Phe Arg Trp Phe Lys Pro Val

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70 1          5
73 <210> SEQ ID NO: 5
74 <211> LENGTH: 8
75 <212> TYPE: PRT
76 <213> ORGANISM: Artificial
78 <220> FEATURE:
79 <223> OTHER INFORMATION: Synthetic derviative of SEQ ID NO:1
82 <220> FEATURE:
83 <221> NAME/KEY: MOD_RES
84 <222> LOCATION: (2)..(2)
85 <223> OTHER INFORMATION: Nle
87 <400> SEQUENCE: 5
W--> 89 His Xaa Arg Trp Phe Lys Pro Val
90 1          5
93 <210> SEQ ID NO: 6
94 <211> LENGTH: 8
95 <212> TYPE: PRT
96 <213> ORGANISM: Artificial
98 <220> FEATURE:
99 <223> OTHER INFORMATION: Synthetic dimer derivative of carboxy terminus of SEQ ID NO:1
102 <220> FEATURE:
103 <221> NAME/KEY: DISULFID
104 <222> LOCATION: (4)..(5)
106 <400> SEQUENCE: 6
108 Val Pro Lys Cys Cys Lys Pro Val
109 1          5
112 <210> SEQ ID NO: 7
113 <211> LENGTH: 3
114 <212> TYPE: PRT
115 <213> ORGANISM: Artificial
117 <220> FEATURE:
118 <223> OTHER INFORMATION: Synthetic derivative of SEQ ID NO:1
121 <220> FEATURE:
122 <221> NAME/KEY: MISC_FEATURE
123 <222> LOCATION: (2)..(2)
124 <223> OTHER INFORMATION: Pro is D-isomer
126 <400> SEQUENCE: 7
128 Lys Pro Val
129 1
132 <210> SEQ ID NO: 8
133 <211> LENGTH: 13
134 <212> TYPE: PRT
135 <213> ORGANISM: Artificial
137 <220> FEATURE:
138 <223> OTHER INFORMATION: Synthetic, modified derivative of SEQ ID NO:1
141 <220> FEATURE:
142 <221> NAME/KEY: MISC_FEATURE
143 <222> LOCATION: (4)..(4)
144 <223> OTHER INFORMATION: X is Nle

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146 <220> FEATURE:
147 <221> NAME/KEY: MISC_FEATURE
148 <222> LOCATION: (7)..(7)
149 <223> OTHER INFORMATION: Phe is D-isomer
151 <400> SEQUENCE: 8
W--> 153 Ser Tyr Ser Xaa Glu His Phe Arg Trp Gly Lys Pro Val
154 1 5 10
157 <210> SEQ ID NO: 9
158 <211> LENGTH: 3
159 <212> TYPE: PRT
160 <213> ORGANISM: Artificial
162 <220> FEATURE:
163 <223> OTHER INFORMATION: Synthetic, modified derivative of SEQ ID NO:1
166 <220> FEATURE:
167 <221> NAME/KEY: MISC_FEATURE
168 <222> LOCATION: (1)..(1)
169 <223> OTHER INFORMATION: Acetylated
171 <220> FEATURE:
172 <221> NAME/KEY: MISC_FEATURE
173 <222> LOCATION: (2)..(2)
174 <223> OTHER INFORMATION: D-isomer of Lys
176 <400> SEQUENCE: 9
178 Lys Pro Val
179 1
182 <210> SEQ ID NO: 10
183 <211> LENGTH: 21
184 <212> TYPE: DNA
185 <213> ORGANISM: Artificial
187 <220> FEATURE:
188 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
190 <400> SEQUENCE: 10
191 gagggcaagc gtcctactc c 21
194 <210> SEQ ID NO: 11
195 <211> LENGTH: 21
196 <212> TYPE: DNA
197 <213> ORGANISM: Artificial
199 <220> FEATURE:
200 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
202 <400> SEQUENCE: 11
203 ggggccctcg tccttcttct c 21
206 <210> SEQ ID NO: 12
207 <211> LENGTH: 19
208 <212> TYPE: DNA
209 <213> ORGANISM: Artificial
211 <220> FEATURE:
212 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
214 <400> SEQUENCE: 12
215 gccaccatcg ccaagaacc 19
218 <210> SEQ ID NO: 13

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219 <211> LENGTH: 19
220 <212> TYPE: DNA
221 <213> ORGANISM: Artificial
223 <220> FEATURE:
224 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
226 <400> SEQUENCE: 13
227 atagccagga agaagacca 19
230 <210> SEQ ID NO: 14
231 <211> LENGTH: 24
232 <212> TYPE: DNA
233 <213> ORGANISM: Artificial
235 <220> FEATURE:
236 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
238 <400> SEQUENCE: 14
239 cggtagccga catgctggta agtg 24
242 <210> SEQ ID NO: 15
243 <211> LENGTH: 24
244 <212> TYPE: DNA
245 <213> ORGANISM: Artificial
247 <220> FEATURE:
248 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
250 <400> SEQUENCE: 15
251 tgaggagcat catggcgaag aaca 24
254 <210> SEQ ID NO: 16
255 <211> LENGTH: 21
256 <212> TYPE: DNA
257 <213> ORGANISM: Artificial
259 <220> FEATURE:
260 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
262 <400> SEQUENCE: 16
263 cattgctgtg gaggtgtttc t 21
266 <210> SEQ ID NO: 17
267 <211> LENGTH: 21
268 <212> TYPE: DNA
269 <213> ORGANISM: Artificial
271 <220> FEATURE:
272 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
274 <400> SEQUENCE: 17
275 gccgtcatga tgtgggtggta g 21
278 <210> SEQ ID NO: 18
279 <211> LENGTH: 26
280 <212> TYPE: DNA
281 <213> ORGANISM: Artificial
283 <220> FEATURE:
284 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
286 <400> SEQUENCE: 18
287 tgaaggtcgg agtcaacgga tttggt 26
290 <210> SEQ ID NO: 19
291 <211> LENGTH: 24

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292 &lt;212&gt; TYPE: DNA

293 &lt;213&gt; ORGANISM: Artificial

295 &lt;220&gt; FEATURE:

296 &lt;223&gt; OTHER INFORMATION: Synthetic oligonucleotide primer

298 &lt;400&gt; SEQUENCE: 19

299 catgtgggcc atgaggtcca ccac

24

## RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 09/07/2006

PATENT APPLICATION: US/10/774,271

TIME: 14:51:05

Input Set : A:\10-774271 - 099425-0100.ST25.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 2

Seq#:8; Xaa Pos. 4

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19

VERIFICATION SUMMARY

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Input Set : A:\10-774271 - 099425-0100.ST25.txt

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L:89 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0